

Fig. 1

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			training										test				discriminant coefficient																			
Symbol	Title	GenBank Accession	node-positive								node-negative								permutational P																	
			104+	75+	72+	63+	61+	59+	58+	51+	48+	101-	97-	73-	57-	47-	106+	105+			137-	138-														
DDOST	dolichyl-diphosphooligosaccharide-protein glycosyltransferase	BF941738	0.83	1.42	2.23	1.32	1.20	0.82	1.60	0.68	1.51	1.29	0.19	1.79	2.94	-0.66	0.57	0.50	1.01	-0.34	-1.09	-0.35	3.5	0.00006	1.87											
GNS	glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	Z12173	1.63	2.56	0.70	1.28	0.56	1.76	0.97	1.93	1.29												0.00119	1.26												
NEDD8	neural precursor cell expressed, developmentally down-regulated 8	D23662	-0.23	-0.10	0.23	0.29	-0.14	1.18	1.16	1.01	0.19												0.00189	1.29												
	DKFZP5661133	AA533633	-0.07	0.09	0.59	0.75	1.33	-0.39	0.80	0.97	1.79												0.00348													
	ESTs, Highly similar to B27079 fibronectin receptor beta chain precursor	A1755112	1.36	1.50	3.65	1.27	2.91	1.31	2.37	1.26	2.94												0.00432													
LOC 51096	CGI-48 protein	AA448503	0.41	-0.14	-0.09	0.69	0.95	0.13	-0.51	0.04	-0.66												0.00677	1.36												
CCT5	chaperonin containing TCP1, subunit 5 (epsilon)	D43950	0.72	0.81	0.60	0.24	0.60	2.03	1.52	2.04	0.57												0.00821													
CCT3	chaperonin containing TCP1, subunit 3 (gamma)	X74801	1.75	1.59	0.24	0.68	0.43	1.24	0.25	1.14	0.50												0.00905													
PPP 2R1B	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR65), beta isoform	M65254	0.36	1.09	0.53	1.15	-0.02	1.60	0.73	1.25	1.01												0.00984													
UBQLN1	ubiquitin 1	AA281115	0.02	-0.14	-1.25	0.29	-0.29	-0.61	-0.13	-0.73	-0.34												0.00002													
AIM2	absent in melanoma 2	AF024714	-0.06	0.18	-0.49	-0.20	-0.45	-0.77	0.10	0.81	-1.09												0.00086	-1.54												
USP9X	ubiquitin specific protease 9, X (chromosome (Drosophila fat facets related))	X98296	-0.13	-0.11	-0.36	0.25	-1.42	-0.43	-1.00	-1.00	-0.35												0.00652													
			"Predictive score"														constant (-1.954)																			
			logarithmic expression ratio										-2.0				-1.6		-1.2		-0.8		-0.4		0.0		0.4		0.8		1.2		1.6		2.0	
			expression ratio(Cy3/Cy5)										0.25				0.33		0.44		0.57		0.76		1.00		1.32		1.74		2.30		3.03		4.00	

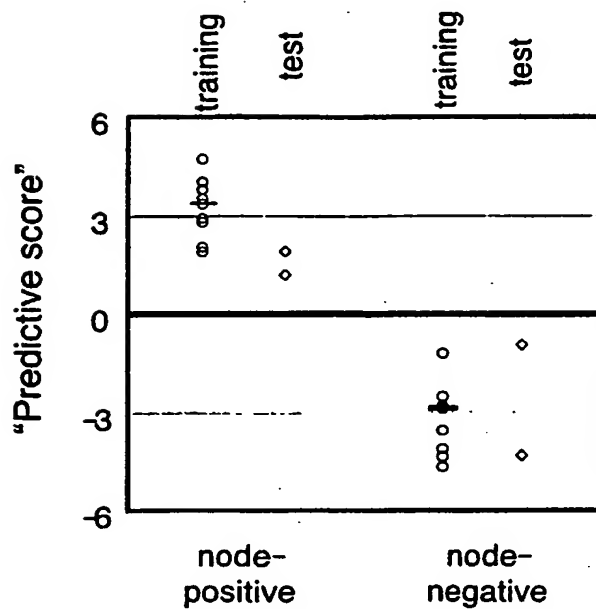


Fig. 2B

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